

Multiscale modelling of sarcoma and bone tissue synergy and interactions

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Multiscale modelling of sarcoma and bone tissue synergy and interactions

Acknowledgments

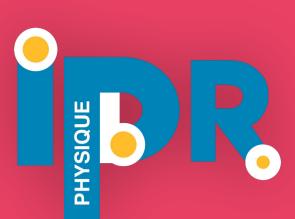
F. Rédini, S. Battaglia

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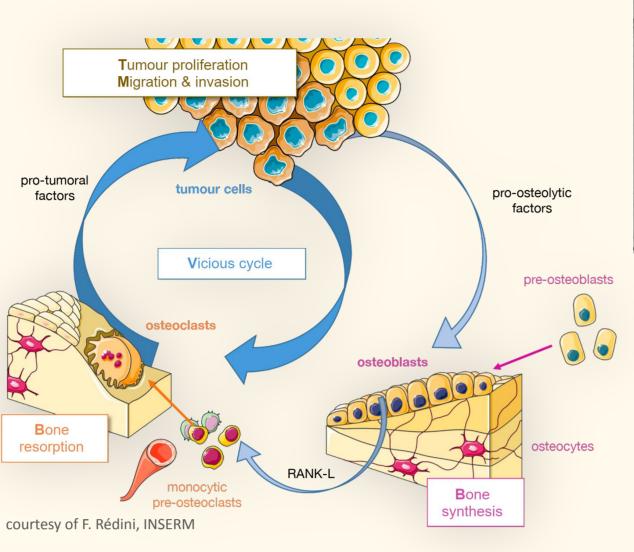


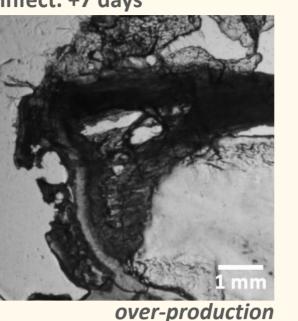
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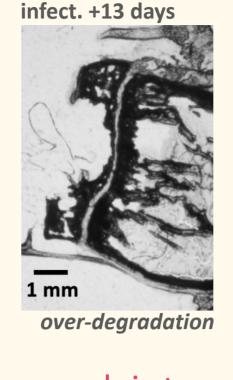
Understanding the morphology of osteosarcoma

Osteosarcoma, along with Ewing sarcoma, is one of the most frequent malignant primary bone tumour, the eighth most common malignancy of childhood.
Osteosarcomas are aggressive tumours, rapidly invasive, which signalling pathways are hardly understood.



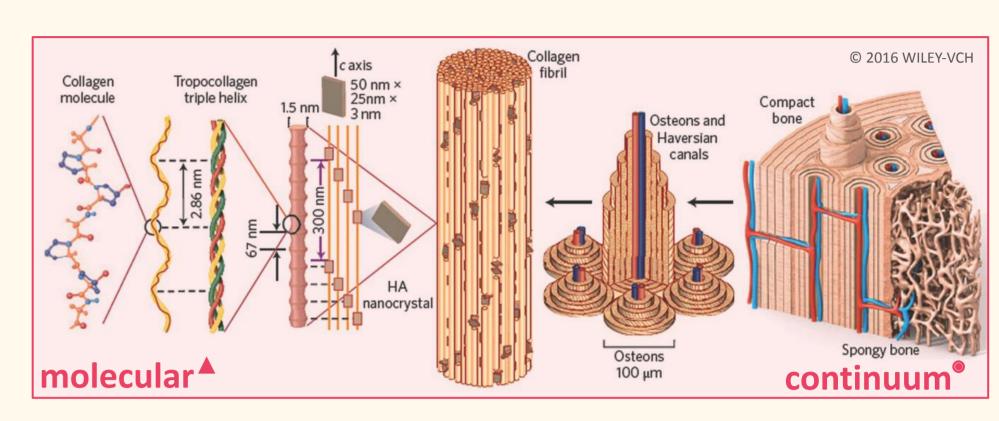






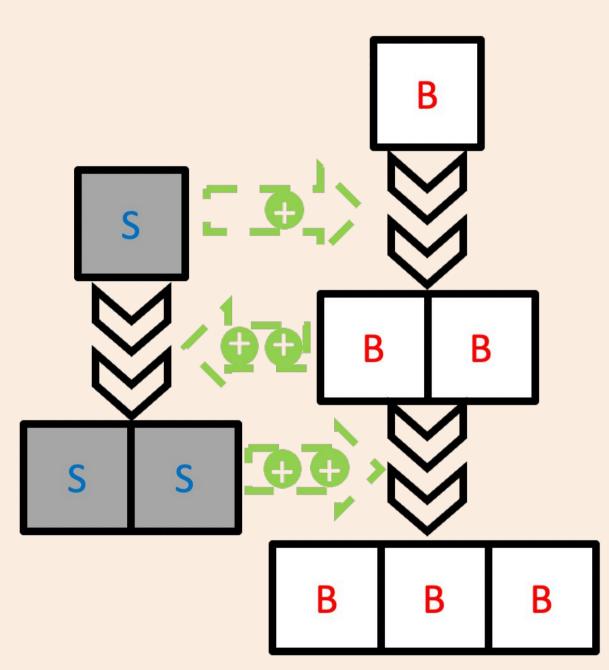
One main area of research is to investigate the inter-dependence of tumour growth and bone remodelling, which constitutes a vicious cycle [1].

The design of treatment or diagnosis requires to clarify the **physical mechanisms** associated with tumour initiation and **growth** at **multiple scales**.



Continuum® modelling of osteosarcomas' growth

On the **microscale**, we simulate bone tissue remodelling and morphology in pathological conditions associated with osteosarcoma using a 3D continuum model based on Cellular Automata (CA).



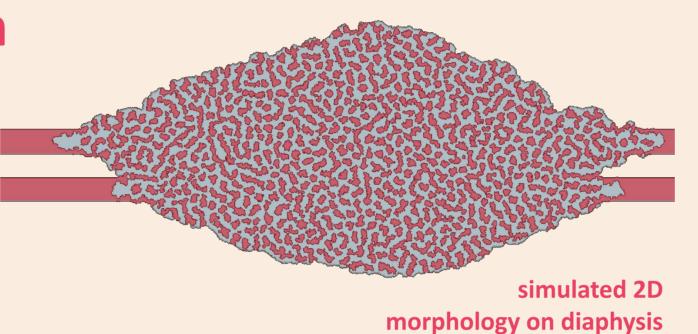
sarcoma and bone synergetic propagation

The model for osteosarcoma and bone morphology growth is formulated based diffusion and reaction mechanisms.

The model accounts for the following mechanisms:

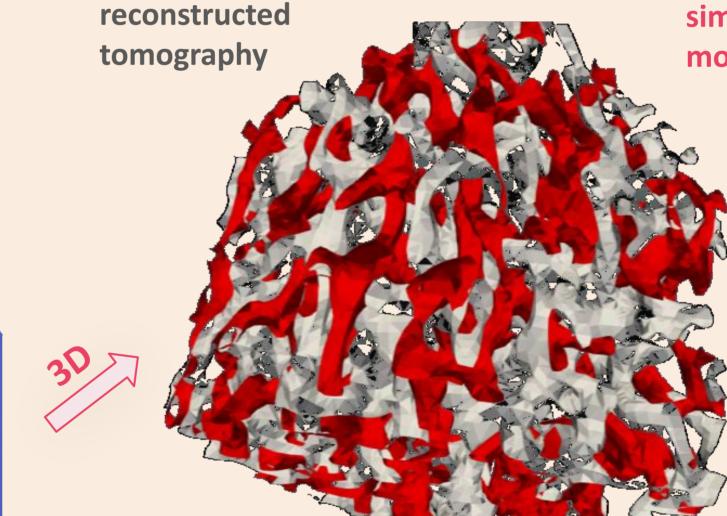
- (i) tumour phase concentration
 promotes bone
 synthesis;
- (ii) bone phase
 concentration
 promotes tumour
 synthesis;
- (iii) the one phase
 synthesis induce the
 other phase resorption;
- (iv) concentration influence is long-range.

The three first mechanisms account for phase production and degradation (reaction), and the last mechanism accounts for signalling agents diffusion.

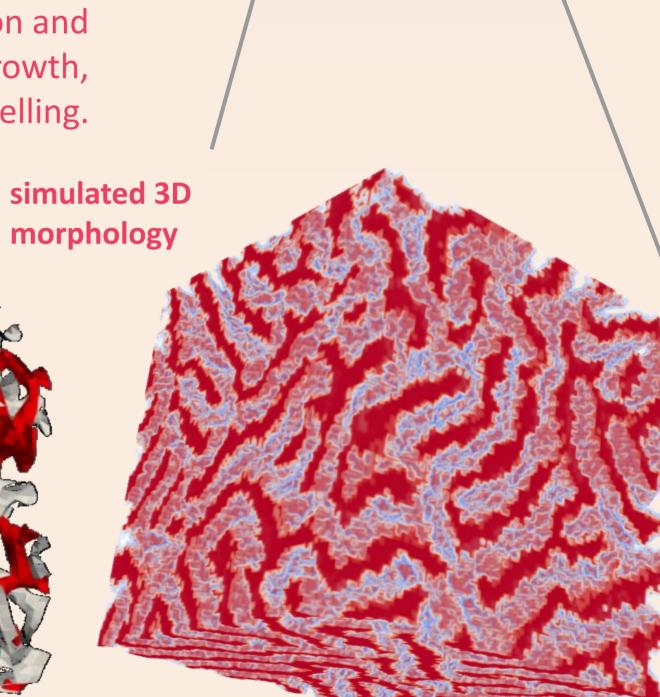


With our simple mechanisms (diffusion, reaction) we **reproduce the** *spongy* **morphology of osteosarcoma**. We can discriminate the simulated morphologies based on **phase connectivity** and **power spectra** with respect to **tomography** data. It is then possible to elucidate characteristic reaction and diffusion timescales in the signalling pathways associated with tumour growth

tomography data. It is then possible to elucidate characteristic reaction and diffusion timescales in the signalling pathways associated with tumour growth, to reveal candidate mediators of the interaction with bone remodelling.

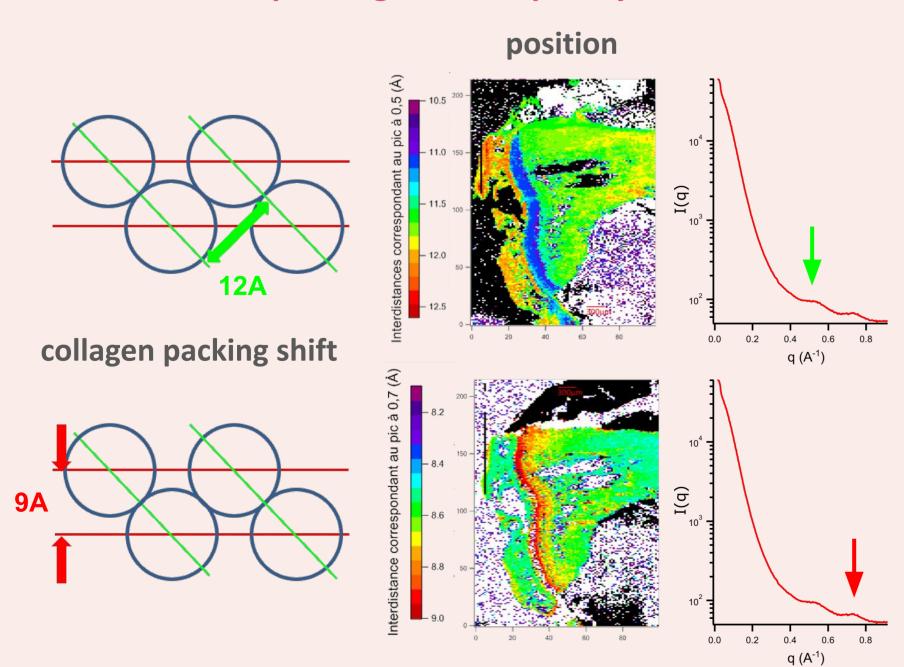


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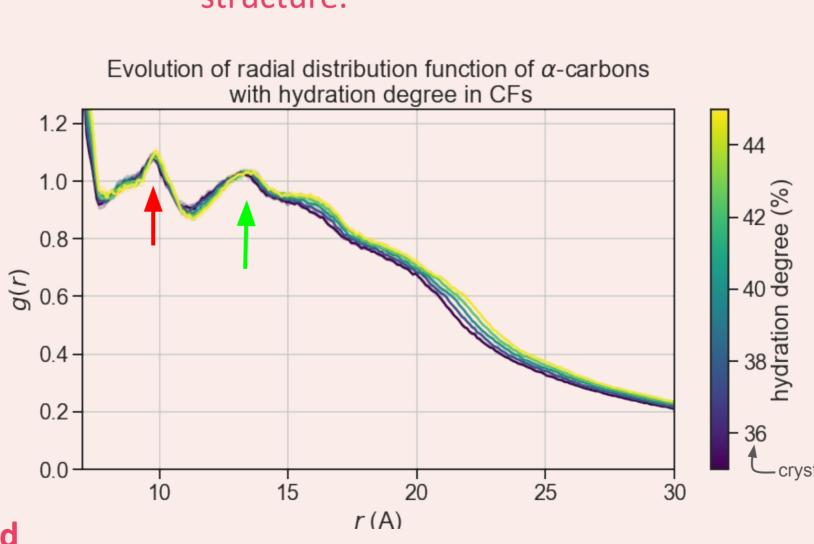


Molecular[▲] modelling of pathological bone tissue self-assembly

We want to explain **early-stage interactions** between primary bone tumours and bone tissue remodelling at the **nanoscale**. We recently correlated **tumour development with rapid changes** in the structure of **Collagen Fibrils** (**CFs**) in bone tissue. However, not only self-organisation but also **structural packing of CFs is poorly elucidated**.



In order to clarify interactions between bone cancer cells and bone tissue remodelling, we designed a nanoscale molecular models of CFs based on X-ray crystallography data [2] and an artificial hydration process, and we simulated hydration up to 50% in water content of the crystallography structure.



We apply the molecular models of microfibril crystals to **understand how rearrangements in CFs could occur**. Molecular dynamics simulations capture the evolution of the **tropocollagen inter-distances** with the degree of hydration or other thermodynamic variables. We then can elucidate the origin of **structural reorganisation associated with sarcomas' growth**.

Simulations of the hydration of the collagen microfibril crystal reveal highly anisotropic water dynamics. We found high water diffusion along the main axis of the tropocollagen molecules, but also in one orthogonal direction. Anisotropy indicate preferential directions for the transport of mineral phase precursors. Our results correlate with the specific platelet morphology of hydroxyapatite crystals within CFs.

References

[1] Corre, I., et al. The osteosarcoma microenvironment: a complex but targetable ecosystem. *Cells* (2020), 9 (4). [2] Orgel, J. P., et al. Microfibrillar structure of type I collagen in situ. *Proc. Natl. Acad. Sci.* (2006) 103 (24).





